PCT/US2004/014540

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SEQUENCE LISTING

:	Isis Pharmaceuticals Inc. Rosanne M. Crooke Mark J. Graham	
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ggt Gl _l	aat Asi	gga 1 Gly 375	, Glr	g agt 1 Ser	tat Tyr	cga Arg	380 380	Thi	tac Tyr	tco Sei	c acc	act Thr 385	· va.	e aca l Thr	a gga Gly	1209
aga	a acc Thi	c Cyr	c caa s Glr	a gct a Ala	tgg Trp	tca Ser 395	: Sei	ato Met	g aca Thi	a cca	a cad o His 400	s sei	g cat	z agt s Sei	cgg Arg	1257
ace Th: 40	r Pro	a gaa o Gli	a tad u Tyi	tac Ty	c cca c Pro 410	a Ası	get n Ala	t ggo a Gl	c tto y Le	ate u Il 41	e Me	g aad t Asi	tac n Ty:	c tgo r Cys	agg Arg 420	1305

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gaa ggg act Glu Gly Thr 2390	gcc gtc gcg Ala Val Ala	cct ccg ac Pro Pro Th 2395	t gtt acc r Val Thr	ccg gtt Pro Val 2400	cca agc Pro Ser	cta 7257 Leu
	tcc gaa caa Ser Glu Gln 241	Ala Pro Th		Arg Pro		
gag tgc tac Glu Cys Tyr	cat ggt aat His Gly Asn 2425	gga cag ag Gly Gln Se	t tat cga r Tyr Arg 2430	ggĉ aca Gly Thr	tac tcc Tyr Ser 2435	Thr
	gga aga acc Gly Arg Thr 2440	Cys Gln Al		Ser Met		
	cgg acc cca Arg Thr Pro 5				Leu Ile	
	agg aat cca Arg Asn Pro					
	ggt gtc agg Gly Val Arg 249	Trp Glu Ty		Leu Thr		
	ggg act gcc	gtc gcg cc Val Ala Pr				

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	2505	2510	2515
agc cta gag gct Ser Leu Glu Ala 2520	Pro Ser Glu Gln	gca ccg act gag of Ala Pro Thr Glu of 2525	caa agg cct ggg 7641 Sln Arg Pro Gly 2530
gtg cag gag tgc Val Gln Glu Cys 2535	tac cat ggt aat Tyr His Gly Asn 254	gga cag agt tat o Gly Gln Ser Tyr i O	ega ggc aca tac 7689 Arg Gly Thr Tyr 2545
tcc acc act gtc Ser Thr Thr Val 2550	aca gga aga acc Thr Gly Arg Thr 2555	tgc caa gct tgg Cys Gln Ala Trp 2560	cca tct atg aca 7737 Ser Ser Met Thr
cca cac tcg cat Pro His Ser His 2565	agt cgg acc cca Ser Arg Thr Pro 2570	gaa tac tac cca Glu Tyr Tyr Pro	aat gct ggc ttg 7785 Asn Ala Gly Leu 2580
atc atg aac tac Ile Met Asn Tyr	tgc agg aat cca Cys Arg Asn Pro 2585	gat gct gtg gca g Asp Ala Val Ala 2 2590	gct cct tat tgt 7833 Ala Pro Tyr Cys 2595
tat acg agg gat Tyr Thr Arg Asp 260	Pro Gly Val Arg	tgg gag tac tgc Trp Glu Tyr Cys 2605	aac ctg acg caa 7881 Asn Leu Thr Gln 2610
tgc tca gac gca Cys Ser Asp Ala 2615	gaa ggg act gcc Glu Gly Thr Ala 262	gtc gcg cct ccg Val Ala Pro Pro 0	act gtt acc ccg 7929 Thr Val Thr Pro 2625
gtt cca agc cta Val Pro Ser Leu 2630	gag gct cct tcc Glu Ala Pro Ser 2635	gaa caa gca ccg Glu Gln Ala Pro 2640	act gag cag agg 7977 Thr Glu Gln Arg
cct ggg gtg cag Pro Gly Val Gln 2645	gag tgc tac cac Glu Cys Tyr His 2650	ggt aat gga cag Gly Asn Gly Gln 2655	agt tat cga ggc 8025 Ser Tyr Arg Gly 2660
aca tac tcc acc Thr Tyr Ser Thr	act gtc act gga Thr Val Thr Gly 2665	aga acc tgc caa Arg Thr Cys Gln 2670	gct tgg tca tct 8073 Ala Trp Ser Ser 2675
	Ser His Ser Arg	acc cca gaa tac Thr Pro Glu Tyr 2685	
		aat cca gat gct Asn Pro Asp Ala 0	
		gtc agg tgg gag Val Arg Trp Glu 2720	
		act gcc gtc gcg Thr Ala Val Ala 2735	

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acc Thr	ccg Pro	gtt Val	cca Pro	agc Ser	cta Leu	gag Glu	gct Ala	cct Pro	Ser	Glu	caa Gln	gca Ala	ccg Pro	Thr	Glu	8313
				2745	5				2750)				2755	1	
caa Gln	agg Arg	cct Pro	999 Gly 2760	Val	cag Gln	gag Glu	tgc Cys	tac Tyr 2765	His	ggt Gly	aat Asn	gga Gly	cag Gln 2770	Ser	tat Tyr	8361
cga Arg	ggc Gly	aca Thr 277	Tyr	tcc Ser	acc Thr	act Thr	gtc Val 2780	Thr	gga Gly	aga Arg	acc Thr	tgc Cys 2789	caa Gln	gct Ala	tgg Trp	8409
tca Ser	tct Ser 2790	Met	aca Thr	cca Pro	cac His	tcg Ser 279	His	agt Ser	cgg Arg	acc Thr	cca Pro 2800	Glu	tac Tyr	tac Tyr	cca Pro	8457
aat Asn 280	Ala	ggc Gly	ttg Leu	atc Ile	atg Met 2810	Asn	tac Tyr	tgc Cys	agg Arg	aat Asn 281	Pro	gat Asp	gct Ala	gtg Val	gca Ala 2820	8505
gct Ala	cct Pro	tat Tyr	tgt Cys	tat Tyr 282	Thr	agg Arg	gat Asp	ccc Pro	ggt Gly 283	Val	agg Arg	tgg Trp	gag Glu	tac Tyr 283	Сув	8553
aac Asn	ctg Leu	acg Thr	caa Gln 284	Cys	tca Ser	gac Asp	gca Ala	gaa Glu 284	Gly	act Thr	gcc Ala	gtc Val	gcg Ala 2850	Pro	ccg Pro	8601
act Thr	gtt Val	acc Thr 285	Pro	gtt Val	cca Pro	agc Ser	cta Leu 286	Glu	gct Ala	cct Pro	tcc Ser	gaa Glu 286	caa Gln 5	gca Ala	ccg Pro	8649
act Thr	gag Glu 287	Gln	agg Arg	cct Pro	gly aaa	gtg Val 287	Gln	gag Glu	tgc Cys	tac Tyr	cat His 288	Gly	aat Asn	gga Gly	cag Gln	8697
agt Ser 288	Tyr	cga Arg	ggc	aca Thr	tac Tyr 289	Ser	acc Thr	act Thr	gtc Val	aca Thr 289	Gly	aga Arg	acc Thr	tgc Cys	caa Gln 2900	8745
gct Ala	tgg Trp	tca Ser	tct Ser	atg Met 290	Thr	cca Pro	cac His	tcg Ser	cat His 291	Ser	cgg Arg	acc Thr	cca Pro	gaa Glu 291	Tyr	8793
tac Tyr	cca Pro	aat Asn	gct Ala 292	Gly	ttg Leu	atc Ile	atg Met	aac Asn 292	\mathtt{Tyr}	tgc Cys	agg Arg	aat Asn	cca Pro 293	Asp	gct Ala	8841
gtg Val	gca Ala	gct Ala 293	Pro	tat Tyr	tgt Cys	tat Tyr	acg Thr 294	Arg	gat Asp	ccc Pro	ggt Gly	gtc Val 294	agg Arg 5	tgg Trp	gag Glu	8889
tac Tyr	tgc Cys 295	Asn	ctg Leu	acg Thr	caa Gln	tgc Cys 295	Ser	gac Asp	gca Ala	gaa Glu	ggg 939	Thr	gcc Ala	gtc Val	gcg Ala	8937
cct	ccg Pro	act Thr	gtt Val	acc Thr	ccg Pro	gtt Val	cca Pro	agc Ser	cta Leu	gag Glu	gct Ala	cct Pro	tcc Ser	gaa Glu	caa Gln	8985

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2965	2970	2975	2980
gca ccg act gag cag Ala Pro Thr Glu Gln 298	Arg Pro Gly Val	g cag gag tgc tac cac L Gln Glu Cys Tyr His 2990	ggt aat 9033 Gly Asn 2995
gga cag agt tat cga Gly Gln Ser Tyr Arg 3000	ggc aca tac tcc Gly Thr Tyr Ser 300	e acc act gtc act gga o Thr Thr Val Thr Gly 05 3010	Arg Thr
tgc caa gct tgg tca Cys Gln Ala Trp Ser 3015	tct atg aca cca Ser Met Thr Pro 3020	a cac tcg cat agt cgg o His Ser His Ser Arg 3025	acc cca 9129 Thr Pro
gaa tac tac cca aat Glu Tyr Tyr Pro Asn 3030	gct ggc ttg ato Ala Gly Leu Ilo 3035	e atg aac tac tgc agg e Met Asn Tyr Cys Arg 3040	aat cca 9177 Asn Pro
gat gct gtg gca gct Asp Ala Val Ala Ala 3045	cct tat tgt tai Pro Tyr Cys Ty: 3050	t acg agg gat ccc ggt Thr Arg Asp Pro Gly 3055	gtc agg 9225 Val Arg 3060
tgg gag tac tgc aac Trp Glu Tyr Cys Asr 306	Leu Thr Gln Cy	c tca gac gca gaa ggg s Ser Asp Ala Glu Gly 3070	act gcc 9273 Thr Ala 3075
gtc gcg cct ccg act Val Ala Pro Pro Thr 3080	gtt acc ccg gt Val Thr Pro Va 30	t cca agc cta gag gct l Pro Ser Leu Glu Ala 35 3090	Pro Ser
		t ggg gtg cag gag tgc o Gly Val Gln Glu Cys 3105	
ggt aat gga cag agt Gly Asn Gly Gln Ser 3110	Tyr Arg Gly Th	a tac tcc acc act gtc r Tyr Ser Thr Thr Val	act gga 9417 Thr Gly
		g aca cca cac tcg cat t Thr Pro His Ser His 3135	
acc cca gaa tac tac Thr Pro Glu Tyr Tyr 314	Pro Asn Ala Gl	c ttg atc atg aac tac y Leu Ile Met Asn Tyr 3150	tgc agg 9513 Cys Arg 3155
aat cca gat gct gtg Asn Pro Asp Ala Val 3160	g gca gct cct ta . Ala Ala Pro Ty 31	t tgt tat acg agg gat r Cys Tyr Thr Arg Asp 65 3170	Pro Gly
gtc agg tgg gag tac Val Arg Trp Glu Tyr 3175	tgc aac ctg ac Cys Asn Leu Th 3180	g caa tgc tca gac gca r Gln Cys Ser Asp Ala 3185	gaa ggg 9609 Glu Gly
act gcc gtc gcg cct Thr Ala Val Ala Pro 3190	ccg act gtt ac Pro Thr Val Th 3195	c ccg gtt cca agc cta r Pro Val Pro Ser Leu 3200	gag gct 9657 Glu Ala

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cct tcc gaa Pro Ser Glu 3205	Gln Ala Pr	g act gag o Thr Glu	cag agg Gln Arg	cct ggg Pro Gly 3215	gtg cag Val Gln	gag tgc Glu Cys 3220	9705
tac cac ggt Tyr His Gly	aat gga ca Asn Gly Gl 3225	ng agt tat n Ser Tyr	cga ggc Arg Gly 323	Thr Tyr	tcc acc Ser Thr	act gtc Thr Val 3235	9753
act gga aga Thr Gly Arg	acc tgc ca Thr Cys Gl 3240	aa gct tgg n Ala Trp	tca tct Ser Ser 3245	atg aca Met Thr	cca cac Pro His 3250	Ser His	9801
agt cgg acc Ser Arg Thr 3255	Pro Glu Ty	ac tac cca yr Tyr Pro 326	Asn Ala	ggc ttg Gly Leu	atc atg Ile Met 3265	aac tac Asn Tyr	9849
tgc agg aat Cys Arg Asn 3270	cca gat go Pro Asp Al	ct gtg gca la Val Ala 3275	gct cct Ala Pro	tat tgt Tyr Cys 3280	Tyr Thr	agg gat Arg Asp	9897
ccc ggt gtc Pro Gly Val 3285	Arg Trp G	ag tac tgc lu Tyr Cys 290	aac ctg Asn Leu	acg caa Thr Gln 3295	tgc tca Cys Ser	gac gca Asp Ala 3300	9945
gaa ggg act Glu Gly Thr	gcc gtc gc Ala Val Al 3305	eg cet ceg la Pro Pro	act gtt Thr Val 331	Thr Pro	gtt cca Val Pro	agc cta Ser Leu 3315	9993
gag gct cct Glu Ala Pro	tcc gaa ca Ser Glu G 3320	aa gca ccg ln Ala Pro	act gag Thr Glu 3325	cag agg Gln Arg	cct ggg Pro Gly 333	Val Gln	10041
gag tgc tac Glu Cys Tyr 333!	His Gly A	at gga cag sn Gly Gln 334	Ser Tyr	cga ggc Arg Gly	aca tac Thr Tyr 3345	tcc acc Ser Thr	10089
act gtc act Thr Val Thr 3350	ggaraga ad Gly Arg T	cc tgc caa hr Cys Gln 3355	rget tgg . Ala Trp	Ser Ser 336	Met Thr	cca cac Pro His	10137
tcg cat agt Ser His Ser 3365	Arg Thr P	ca gaa tac ro Glu Tyr 370	tac cca Tyr Pro	aat gct Asn Ala 3375	ggc ttg Gly Leu	atc atg Ile Met 3380	10185
aac tac tgc Asn Tyr Cys	agg aat c Arg Asn P 3385	ca gat cct ro Asp Pro	gtg gca Val Ala 339	Ala Pro	tat tgt Tyr Cys	tat acg Tyr Thr 3395	10233
agg gat ccc Arg Asp Pro	agt gtc a Ser Val A 3400	gg tgg gag rg Trp Glu	tac tgc Tyr Cys 3405	aac ctg Asn Leu	aca caa Thr Gln 341	Cys Ser	10281
gac gca gaa Asp Ala Glu 341	Gly Thr A	cc gtc gcg la Val Ala 342	Pro Pro	act att Thr Ile	acc ccg Thr Pro 3425	att cca Ile Pro	10329
agc cta gag Ser Leu Glu	gct cct t	ct gaa caa	gca cca	act gag	caa agg	cct ggg	10377

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3430	3435	3440
gtg cag gag tgc tac cac Val Gln Glu Cys Tyr His 3445 345	Gly Asn Gly Gln Ser	Tyr Gln Gly Thr Tyr
ttc att act gtc aca gga Phe Ile Thr Val Thr Gly 3465	aga acc tgc caa gct Arg Thr Cys Gln Ala 3470	tgg tca tct atg aca 10473 Trp Ser Ser Met Thr 3475
cca cac tcg cat agt cgg Pro His Ser His Ser Arg 3480	acc cca gca tac tac Thr Pro Ala Tyr Tyr 3485	cca aat gct ggc ttg 10521 Pro Asn Ala Gly Leu 3490
atc aag aac tac tgc cga Ile Lys Asn Tyr Cys Arg 3495	aat cca gat cct gtg Asn Pro Asp Pro Val 3500	gca gcc cct tgg tgt 10569 Ala Ala Pro Trp Cys 3505
tat aca aca gat ccc agt Tyr Thr Thr Asp Pro Ser 3510	gtc agg tgg gag tac Val Arg Trp Glu Tyr 3515	tgc aac ctg aca cga 10617 Cys Asn Leu Thr Arg 3520
tgc tca gat gca gaa tgg Cys Ser Asp Ala Glu Trp 3525 353	Thr Ala Phe Val Pro	Pro Asn Val Ile Leu
gct cca agc cta gag gct Ala Pro Ser Leu Glu Ala 3545		
ccc ggg gta cag gac tgc Pro Gly Val Gln Asp Cys 3560	tac tac cat tat gga Tyr Tyr His Tyr Gly 3565	cag agt tac cga ggc 10761 Gln Ser Tyr Arg Gly 3570
aca tac tcc acc act gtc Thr Tyr Ser Thr Thr Val 3575		
atg aca cca cac cag cat Met Thr Pro His Gln His 3590	agt cgg acc cca gaa Ser Arg Thr Pro Glu 3595	aac tac cca aat gct 10857 Asn Tyr Pro Asn Ala 3600
ggc ctg acc agg aac tac Gly Leu Thr Arg Asn Tyr 3605 361	Cys Arg Asn Pro Asp	Ala Glu Ile Arg Pro
tgg tgt tac acc atg gat Trp Cys Tyr Thr Met Asp 3625		
aca caa tgc ctg gtg aca Thr Gln Cys Leu Val Thr 3640		
gtc cca gat cca agc aca Val Pro Asp Pro Ser Thr 3655	gag gct tct tct gaa Glu Ala Ser Ser Glu 3660	gaa gca cca acg gag 11049 Glu Ala Pro Thr Glu 3665

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caa Gln	agc Ser 3670	Pro	Gly aaa	gtc Val	cag Gln	gat Asp 3675	Cys	tac Tyr	cat His	ggt Gly	gat Asp 3680	Gly	cag Gln	agt Ser	tat Tyr	11097
cga Arg 3685	ggc Gly	tca Ser	ttc Phe	tct Ser	acc Thr 3690	Thr	gtc Val	aca Thr	gga Gly	agg Arg 3695	Thr	tgt Cys	cag Gln	tct Ser	tgg Trp 3700	11145
tcc Ser	tct Ser	atg Met	aca Thr	cca Pro 3705	His	tgg Trp	cat His	cag Gln	agg Arg 3710	Thr	aca Thr	gaa Glu	tat Tyr	tat Tyr 3715	Pro	11193
aat Asn	ggt Gly	ggc	ctg Leu 3720	Thr	agg Arg	aac Asn	tac Tyr	tgc Cys 3725	Arg	aat Asn	cca Pro	gat Asp	gct Ala 3730	Glu	att Ile	11241
agt Ser	cct Pro	tgg Trp 3735	Cys	tat Tyr	acc Thr	atg Met	gat Asp 3740	Pro	aat Asn	gtc Val	aga Arg	tgg Trp 3745	Glu	tac Tyr	tgc Cys	11289
aac Asn	ctg Leu 3750	Thr	caa Gln	tgt Cys	cca Pro	gtg Val 375	Thr	gaa Glu	tca Ser	agt Ser	gtc Val 3760	Leu	gcg Ala	acg Thr	tcc Ser	11337
acg Thr 376	gct Ala	gtt Val	tct Ser	gaa Glu	caa Gln 3770	Ala	cca Pro	acg Thr	gag Glu	caa Gln 3775	Ser	ccc Pro	aca Thr	gtc Val	cag Gln 3780	11385
gac Asp	tgc Cys	tac Tyr	cat His	ggt Gly 378	Asp	gga Gly	cag Gln	agt Ser	tat Tyr 3790	Arg	ggc Gly	tca Ser	ttc Phe	tcc Ser 379!	Thr	11433
act Thr	gtt Val	aca Thr	gga Gly 380	Arg	aca Thr	tgt Cys	cag Gln	tct Ser 380	Trp	tcc Ser	tct Ser	atg Met	aca Thr 381	Pro	cac His	11481
tgg Trp	cat His	cag Gln 381	Arg	acc Thr	aca Thr	gáa Glu	tac Tyr 382	Tyr	cca Pro	aat Asn	ggt Gly	ggč Gly 382!	Leu	acc Thr	agg Arg	11529
aac Asn	tac Tyr 383	Cys	agg Arg	aat Asn	cca Pro	gat Asp 383	Ala	gag Glu	att Ile	cgc Arg	cct Pro 384	Trp	tgt Cys	tat Tyr	acc Thr	11577
atg Met 384	gat Asp 5	ccc Pro	agt Ser	gtc Val	aga Arg 385	Trp	gag Glu	tac Tyr	tgc Cys	aac Asn 385	Leu	acg Thr	caa Gln	tgt Cys	cca Pro 3860	11625
gtg Val	atg Met	gaa Glu	tca Ser	act Thr 386	Leu	ctc Leu	aca Thr	act Thr	ccc Pro 387	Thr	gtg Val	gtc Val	cca Pro	gtt Val 387	Pro	11673
agc Ser	aca Thr	gag Glu	ctt Leu 388	Pro	tct Ser	gaa Glu	gaa Glu	gca Ala 388	Pro	act Thr	gaa Glu	aac Asn	agc Ser 389	Thr	gly aaa	11721
gtc Val	cag Gln	gac Asp	tgc Cys	tac Tyr	cga Arg	ggt Gly	gat Asp	gga Gly	cag Gln	agt Ser	tat Tyr	cga Arg	ggc	aca Thr	ctc Leu	11769

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3895		3900	3905	
tcc acc act a Ser Thr Thr I 3910	tc aca gga aga le Thr Gly Arg 3915	Thr Cys Gln Ser	egg teg tet atg aca Trp Ser Ser Met Thr 3920	11817
cca cat tgg c Pro His Trp H 3925	eat cgg agg atc His Arg Arg Ile 3930	cca tta tac tat of Pro Leu Tyr Tyr 1	cca aat gct ggc ctg Pro Asn Ala Gly Leu 3940	11865
acc agg aac t Thr Arg Asn T	ac tgc agg aat Yr Cys Arg Asn 3945	cca gat gct gag a Pro Asp Ala Glu : 3950	att cgc cct tgg tgt Ile Arg Pro Trp Cys 3955	11913
Tyr Thr Met A	gat ccc agt gtc Asp Pro Ser Val 3960	agg tgg gag tac Arg Trp Glu Tyr (3965	tgc aac ctg aca cga Cys Asn Leu Thr Arg 3970	11961
tgt cca gtg a Cys Pro Val T 3975	aca gaa tcg agt Thr Glu Ser Ser	gtc ctc aca act Val Leu Thr Thr 3980	ccc aca gtg gcc ccg Pro Thr Val Ala Pro 3985	12009
gtt cca agc a Val Pro Ser 1 3990	aca gag gct cct Thr Glu Ala Pro 3999	Ser Glu Gln Ala	cca cct gag aaa agc Pro Pro Glu Lys Ser 4000	12057
cct gtg gtc o Pro Val Val o 4005	cag gat tgc tac Gln Asp Cys Tyr 4010	cat ggt gat gga His Gly Asp Gly 4015	cgg agt tat cga ggc Arg Ser Tyr Arg Gly 4020	12105
ata tcc tcc a	acc act gtc aca Thr Thr Val Thr 4025	gga agg acc tgt Gly Arg Thr Cys 4030	caa tct tgg tca tct Gln Ser Trp Ser Ser 4035	12153
Met Ile Pro H	cac tgg cat cag His Trp His Gln 4040	agg acc cca gaa Arg Thr Pro Glu 4045	aac tac cca aat gct Asn Tyr Pro Asn Ala 4050	12201
ggc ctg acc of Gly Leu Thr (gag aac tac tgc Glu Asn Tyr Cys	agg aat cca gat Arg Asn Pro Asp 4060	tct ggg aaa caa ccc Ser Gly Lys Gln Pro 4065	12249
tgg tgt tac a Trp Cys Tyr 1 4070	aca acc gat ccg Thr Thr Asp Pro 407	Cys Val Arg Trp	gag tac tgc aat ctg Glu Tyr Cys Asn Leu 4080	12297
aca caa tgc Thr Gln Cys 4085	tca gaa aca gaa Ser Glu Thr Glu 4090	tca ggt gtc cta Ser Gly Val Leu 4095	gag act ccc act gtt Glu Thr Pro Thr Val 4100	12345
gtt cca gtt Val Pro Val	cca agc atg gag Pro Ser Met Glu 4105	gct cat tct gaa Ala His Ser Glu 4110	gca gca cca act gag Ala Ala Pro Thr Glu 4115	12393
Gln Thr Pro	gtg gtc cgg cag Val Val Arg Gln 4120	tgc tac cat ggt Cys Tyr His Gly 4125	aat ggc cag agt tat Asn Gly Gln Ser Tyr 4130	12441

cga Arg	ggc Gly	aca Thr 4135	Phe	tcc Ser	acc Thr	act Thr	gtc Val 4140	Thr	gga Gly	agg Arg	aca Thr	tgt Cys 4145	Gln	tct Ser	tgg Trp	12489
tca Ser	tcc Ser 4150	Met	aca Thr	cca Pro	cac His	cgg Arg 4155	His	cag Gln	agg Arg	acc Thr	cca Pro 4160	Glu	aac Asn	tac Tyr	cca Pro	12537
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atg Met	ttt Phe 423	Gly	aat Asn	Gly 999	aaa Lys	gga Gly 423	Tyr	cgg Arg	ggc Gly	aag Lys	aag Lys 424	Ala	acc Thr	act Thr	gtt Val	12777
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agc Ser	acg Thr	ttc Phe	att Ile	cca Pro 426	Gly	aca Thr	aat Asn	aaa Lys	tgg Trp 427	Ala	ggt Gly	ctg Leu	gaa Glu	aaa Lys 427	Asn	12873
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atg Met	aat Asn	cca Pro 429	Arg	aaa Lys	ctt Leu	ttt Phe	gac Asp 430	\mathtt{Tyr}	tgt Cys	gat Asp	atc Ile	Pro 430	Leu	tgt Cys	gca Ala	12969
tcc Ser	tct Ser 431	Ser	ttt Phe	gat Asp	tgt Cys	999 Gly 431	Lys	cct Pro	caa Gln	gtg Val	gag Glu 432	Pro	aag Lys	aaa Lys	tgt. Cys	13017
cct Pro	gga	agc Ser	att Ile	gta Val	ggg Gly	Gly	tgt Cys	gtg Val	gcc Ala	cac His 433	Pro	cat His	tcc Ser	tgg Trp	ccc Pro 4340	13065
432	_	552			433	0				433	_					
taa	_	atc	aqt	ctc	aga Arg	aca	agg Arg	ttt Phe	gga Gly 435	aag Lys	cac	ttc Phe	tgt Cys	gga Gly 435	ggc Gly	13113

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gaa gtg aac ct Glu Val Asn Le 4390	c gaa tct cat gtt u Glu Ser His Val 4395	cag gaa ata gaa Gln Glu Ile Glu 4400	Val Ser Arg Leu	13257
ttc ttg gag cc Phe Leu Glu Pro 4405	c aca caa gca gat o Thr Gln Ala Asp 4410	att gcc ttg cta Ile Ala Leu Leu 4415	aag cta agc agg Lys Leu Ser Arg 4420	13305
cct gcc gtc at Pro Ala Val Il	c act gac aaa gta e Thr Asp Lys Val 4425	atg cca gct tgt Met Pro Ala Cys 4430	ctg cca tcc cca Leu Pro Ser Pro 4435	13353
Asp Tyr Met Va	c acc gcc agg act l Thr Ala Arg Thr 40	gaa tgt tac atc Glu Cys Tyr Ile 4445	act ggc tgg gga Thr Gly Trp Gly 4450	13401
gaa acc caa gg Glu Thr Gln Gl 4455	t acc ttt ggg act y Thr Phe Gly Thr 446	Gly Leu Leu Lys	gaa gcc cag ctc Glu Ala Gln Leu 4465	13449
ctt gtt att ga Leu Val Ile Gl 4470	ng aat gaa gtg tgc nu Asn Glu Val Cys 4475	aat cac tat aag Asn His Tyr Lys 4480	Tyr Ile Cys Ala	13497
gag cat ttg gc Glu His Leu Al 4485	cc aga ggc act gac .a Arg Gly Thr Asp 4490	agt tgc cag ggt Ser Cys Gln Gly 4495	gac agt gga ggg Asp Ser Gly Gly 4500	13545
cct ctg gtt tg Pro Leu Val Cy	gc ttc gag aag gac rs Phe Glu Lys Asp 4505	aaa tac att tta Lys Tyr Ile Leu 4510	caa gga gtc act Gln Gly Val Thr 4515	13593
Ser Trp Gly Le	et ggc tgt gca cgc eu Gly Cys Ala Arg 520	ccc aat aag cct Pro Asn Lys Pro 4525	ggt gtc tat gct Gly Val Tyr Ala 4530	13641
cgt gtt tca ag Arg Val Ser Ar 4535	gg ttt gtt act tgg rg Phe Val Thr Trp 454	Ile Glu Gly Met	atg aga aat aat Met Arg Asn Asn 4545	13689
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<400> 7 cccggtgtca ggtgggagta ctgc	24
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<220> <223> PCR Primer	
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